

George G. Vega Yon, Ph.D.

mo: (626) 381-8171 | email: g.vegayon@gmail.com | Sandy, UT, 84094

<https://ggv.cl> / <https://github.com/gvegayon> / <https://linkedin.com/in/georgevegayon> / <https://ggv.cl/scholar>

PROFESSIONAL SUMMARY

Accomplished **data scientist** with over **10 years of experience** with multiple software packages and scientific publications. A results-driven professional with over **half a million downloads** and over **200 citations for academic work**. A solution development and delivery champion with a proven track record of spearheading solutions across data science, network science, and statistics.

EXPERIENCE

Research Assistant Professor in Epidemiology, The University of Utah

Nov 2021 – Present

- Studies Machine Learning, Network Science, Data Science, Statistical Computing.
- Managed a team of researchers, leading to published papers, software, and conference talks.
- Taught and designed the first course on HPC using R and C++ (graduate level).
- Founder of the "Network Science and Social Network Analysis at the U" (NetSNAU) Group.
- Contributed to research grants (CDC and VA,) helping to secure over 1 MM USD in funding.
- Core faculty member of the "Utah Center for Data Science".

Research Programmer II, University of Southern California

Feb 2018 – Nov 2021

- Provides technical support and training sessions on software development, HPC, R, and C++.
- Write scientific papers on network science, statistics, and phylogenetics and present them at conferences.
- Designed and taught the course "Intro to Health Data Science" (graduate level).
- Contributed to research grants (NIH and DoD,) helping to secure over 10 MM USD in funding.

Programmer Analyst II, University of Southern California

Oct 2015 – Feb 2018

- Organized local conferences on Network Science.
- Founder of the "R Bookcamp for Statistical Computing."
- Wrote scientific papers and software on network science and presented them at conferences.
- Designed and led workshops on R and Social Network Analysis.

Research Analyst, Chilean Pension Supervisor

Aug 2011 – Aug 2014

- Wrote papers and automatized statistical reports about the Chilean unemployment insurance system.
- Managed social security records and created representative samples for researchers.
- Designed and implemented a pipeline for simulation and forecasting of the unemployment insurance government funds. Reports were distributed to the Chilean Congress.

EDUCATION

Ph.D. in Biostatistics

University of Southern California, 2020

M.Sc. in Economics

California Institute of Technology, 2016

MA in Economics and Public Policy

Universidad Adolfo Ibáñez, 2011

BS in Business Administration

Universidad Adolfo Ibáñez, 2011

SKILLS

R, C++, LaTeX, SQL, Python, XML, NLP, Stata, AWS, Git, GitHub, Docker, tensorflow, continuous integration, Slurm, Unix, Jira, Scrum, Kanban, team management and coordination, R Shiny, excellent communication skills.

SOFTWARE PACKAGES (selected)

- *aphylo: Statistical Inference of Annotated Phylogenetic Trees* (2022). R package version 0.2-1 URL: <https://cran.r-project.org/package=aphylo> .
- *rgexf: Build, Import and Export GEXF Graph Files* (2020). R package version 0.16.0. URL: <https://CRAN.R-project.org/package=rgexf>.
- *netdiffuseR: Analysis of Diffusion and Contagion Processes on Networks* (2020). R package version 1.22.0. URL: <https://cran.r-project.org/package=netdiffuseR>.
- *ergmito: Exponential Random Graph Models for Small Networks* (2020). R package version 0.3-0. URL: <https://cran.r-project.org/package=ergmito>.
- *slurmR: A Lightweight Wrapper for 'Slurm'* (2020). R package version 0.4-1. URL: <https://CRAN.R-project.org/package=slurmR>.
- *fmcmmc: A friendly MCMC framework* (2020). R package version 0.3-0. URL: <https://CRAN.R-project.org/package=fmcmmc> .

ACADEMIC PUBLICATIONS (selected)

- George G. Vega Yon. “Power and Multicollinearity in Small Networks: A Discussion of “Tale of Two Datasets: Representativeness and Generalisability of Inference for Samples of Networks” by Krivitsky, Coletti & Hens”. In: **Journal of The American Statistical Association** (2023). to appear.
- George G. Vega Yon, Mary Jo Pugh, and Thomas W. Valente. *Discrete Exponential-Family Models for Multivariate Binary Outcomes*. Nov. 2022. In: **arXiv**: [2211.00627](https://arxiv.org/abs/2211.00627) [cs. stat].
- George G. Vega Yon, Andrew Slaughter, and Kayla de la Haye. “Exponential random graph models for little networks”. In: **Social Networks** 64 (2021), pp. 225–238. URL: <https://doi.org/10.1016/j.socnet.2020.07.005>.
- George G. Vega Yon, Duncan C. Thomas, John Morrison, Huaiyu Mi, et al. “Bayesian parameter estimation for automatic annotation of gene functions using observational data and phylogenetic trees”. In: **PLOS Computational Biology** 17.2 (Feb. 2021), pp. 1–35. URL: <https://doi.org/10.1371/journal.pcbi.1007948>.
- George G. Vega Yon and Paul Marjoram. “fmcmmc: A friendly MCMC framework”. In: **Journal of Open Source Software** 4.39 (July 2019), p. 1427. URL: <http://joss.theoj.org/papers/10.21105/joss.01427>.
- George G. Vega Yon and Brian Quistorff. “parallel: A command for parallel computing”. In: **The Stata Journal: Promoting communications on statistics and Stata** 19.3 (Sept. 2019), pp. 667–684. URL: <http://journals.sagepub.com/doi/10.1177/1536867X19874242>.

AWARDS

- Best Paper Awards, 72 ICA conference, 2022.
- Travel Grant, Society of Young Network Scientist, 2019.
- Fellowship, California Institute of Technology, 2014.
- Scholarship, Adolfo Ibáñez University, 2006.

REFEREE (ad hoc reviewer)

- Journal of The American Statistical Association
- BMC Infectious Diseases
- The Official Journal of The Society for Computational Economics
- The R Journal
- Social Networks
- Journal of Mathematical Sociology
- Journal of Open Source Software
- Bioinformatics
- Computer Methods and Programs in Biomedicine Update